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- AN - 1996:515033 BIOSIS
- DN - PREV199699237389
- TI - Analysis of hardening related proteins in *Lolium temulentum* L.
- IN - Tase, Kazuhiro (1); Kobayashi, Makoto; Fujii, Hiroki (1)
- CS - (1) Hokuriku Natl. Agric. Experiment Stn., Inada, Joetsu, Niigata 943-01 Japan
- SO - Grassland Science, (1996) Vol. 42, No. 2, pp. 117-122. ISSN: 0447-5933.
- DT - Article
- LA - English
- SL - English; Japanese
- AB - This study was carried out to identify protein changes occurring through hardening in *Lolium temulentum* L. and to analyze N-terminal amino acid sequences of these hardening related proteins. The development of the freezing tolerance was mostly obtained after 2 weeks exposure to hardening treatment. Thus, the proteins extracted from seedlings of non-hardened and hardened for 2 weeks were analyzed by two-dimensional polyacrylamide gel electrophoresis (2DPAGE). One newly induced protein and six proteins increased in intensity were identified in hardened seedlings, as compared to non-hardened seedlings. To get some information about these proteins, N-terminal amino acid sequences analysis were carried out using gas-phase protein sequencer and N-terminal amino acid sequences of five proteins were determined. As the result of homology search with a protein sequence data bank, several of these proteins were homologous to abscisic acid (ABA)-inducible protein of alfalfa (*Medicago sativa* L.), ribulose-bisphosphate carboxylase (Rubisco) large-subunit binding protein beta chain in garden pea (*Pisum sativum* L.) and leaf fructose diphosphate aldolase in spinach (*Spinacia oleracea* L.). The information of partial amino acid sequences would be helpful in estimating the function of hardening related proteins, and for molecular cloning.
- CC - External Effects - Temperature as a Primary Variable - Cold *10616
Metabolism - Proteins, Peptides and Amino Acids *13012
Temperature: Its Measurement, Effects and Regulation -
Thermoadaptation *23010
Plant Physiology, Biochemistry and Biophysics - Metabolism *51519
Plant Physiology, Biochemistry and Biophysics - Chemical
Constituents *51522
- BC - Gramineae 25305
Chenopodiaceae 25795
Leguminosae *26260
- IT - Major Concepts
Biochemistry and Molecular Biophysics; Metabolism; Physiology
- IT - Chemicals & Biochemicals
POLYACRYLAMIDE; ABSCISIC ACID; RIBULOSE-1,5-BISPHOSPHATE
CARBOXYLASE-OXYGENASE; FRUCTOSE DIPHOSPHATE ALDOLASE
- IT - Sequence Data
amino acid sequence; molecular sequence data
- IT - Miscellaneous Descriptors
ABSCISIC ACID; ANALYTICAL METHOD; BIOCHEMISTRY AND BIOPHYSICS;
CHEMICAL COORDINATION; FREEZING TOLERANCE;
FRUCTOSE DIPHOSPHATE ALDOLASE; METABOLISM; MOLECULAR CLONING
USEFULNESS; POLYACRYLAMIDE GEL ELECTROPHORESIS;
RIBULOSE-1,5-BISPHOSPHATE CARBOXYLASE-OXYGENASE
- ORGN- Super Taxa
Chenopodiaceae: Dicotyledones, Angiospermae, Spermatophyta,

Plantae; Gramineae: Monocotyledones, Angiospermae, Spermatophyta,
Plantae; Leguminosae: Dicotyledones, Angiospermae, Spermatophyta,
Plantae; Plantae - Unspecified: Plantae

ORGN- Organism Name

alfalfa (Leguminosae); garden pea (Leguminosae); plant (Plantae -
Unspecified); spinach (Chenopodiaceae); Lolium
temulentum (Gramineae); Medicago sativa (Leguminosae); Pisum
sativum (Leguminosae); Spinacia oleracea (Chenopodiaceae)

ORGN- Organism Superterms

angiosperms; dicots; monocots; plants; spermatophytes; vascular
plants

RN - 9003-05-8 (POLYACRYLAMIDE)

21293-29-8 (ABSCISIC ACID)

9027-23-0 (RIBULOSE-1,5-BISPHOSPHATE CARBOXYLASE-OXYGENASE)

9024-52-6 (FRUCTOSE DIPHOSPHATE ALDOLASE)